SEQUENCE LISTING

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<NO> Evans, Ronald M.
Chen, J. Don
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<120> A FAMILY OF TRANSCRIPTIONAL CO-REPRESSORS THAT INTERACT WITH NUCLEAR HORMONE RECEPTORS AND USES THEREFOR

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Ser Asn Pro Pro Leu Ile Ser Ser Ala Lys His Pro Ser Val Leu
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Val Pro Tyr Ser Glu His Ala Lys Ala Pro Val Gly Pro Val Thr Met
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Thr Arg Val Pro Ser Asp Ser Ala Ile Thr Tyr Arg Gly Ser Ile Thr

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His Gly Thr Pro Ala Asp Val Leu Tyr Lys Gly Thr lie Thr Arg lie 210 215 220

Ile Gly Glu Asp Ser Pro Ser Arg Leu Asp Arg Gly Arg Glu Asp Ser 225 230 235 240

Leu Pro Lys Gly His Val Ile Tyr Glu Gly Lys Lys Gly His Val Leu 245 250 255

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Ala Tyr Leu Pro Thr Ala Pro Gln Pro Phe Ser Ser Arg His Ser Ser 745 Ser Pro Leu Ser Pro Gly Gly Pro Thr His Leu Thr Lys Pro Thr Thr 760 Thr Ser Ser Ser Glu Arg Glu Arg Asp Arg Asp Arg Glu Arg Asp Arg 775 780 Asp Arg Glu Arg Glu Lys Ser Ile Leu Thr Ser Thr Thr Thr Val Glu 795 790 His Ala Pro Ile Trp Arg Pro Gly Thr Glu Gln Ser Ser Gly Ser Ser 810 805 Gly Ser Ser Gly Gly Gly Gly Ser Ser Arg Pro Ala Ser His 825 820 Ser His Ala His Gln His Ser Pro Ile Ser Pro Arg Thr Gln Asp Ala 840 Leu Gln Gln Arq Pro Ser Val Leu His Asn Thr Gly Met Lys Gly Ile 855 860 Ile Thr Ala Val Glu Pro Ser Lys Pro Thr Val Leu Arg Ser Thr Ser 875 870 Thr Ser Ser Pro Val Arg Pro Ala Ala Thr Phe Pro Pro Ala Thr His 890 885 Cys Pro Leu Gly Gly Thr Leu Asp Gly Val Tyr Pro Thr Leu Met Glu 905 Pro Val Leu Leu Pro Lys Glu Ala Pro Arg Val Ala Arg Pro Glu Arg 920 Pro Arg Ala Asp Thr Gly His Ala Phe Leu Ala Lys Pro Pro Ala Arg 935 940 Ser Gly Leu Glu Pro Ala Ser Ser Pro Ser Lys Gly Ser Glu Pro Arg 950 955 Pro Leu Val Pro Pro Val Ser Gly His Ala Thr Ile Ala Arg Thr Pro 970 965 Ala Lys Asn Leu Ala Pro His His Ala Ser Pro Asp Pro Pro Ala Pro 980 985 990 Pro Ala Ser Ala Ser Asp Pro His Arg Glu Lys Thr Gln Ser Lys Pro 1005 1000 Phe Ser Ile Gln Glu Leu Glu Leu Arg Ser Leu Gly Tyr His Gly Ser 1015 1020 Ser Tyr Ser Pro Glu Gly Val Glu Pro Val Ser Pro Val Ser Ser Pro 1035 1030 Ser Leu Thr His Asp Lys Gly Leu Pro Lys His Leu Glu Glu Leu Asp 1050 1045 Lys Ser His Leu Glu Gly Glu Leu Arg Pro Lys Gln Pro Gly Pro Val 1065 1070 Lys Leu Gly Gly Glu Ala Ala His Leu Pro His Leu Arg Pro Leu Pro 1080 1085 Glu Ser Gln Pro Ser Ser Ser Pro Leu Leu Gln Thr Ala Pro Gly Val 1095 1100 Lys Gly His Gln Arg Val Val Thr Leu Ala Gln His Ile Ser Glu Val 1110 1115 Ile Thr Gln Asp Tyr Thr Arg His His Pro Gln Gln Leu Ser Ala Pro 1125 1130 Leu Pro Ala Pro Leu Tyr Ser Phe Pro Gly Ala Ser Cys Pro Val Leu 1145 1140 Asp Leu Arg Arg Pro Pro Ser Asp Leu Tyr Leu Pro Pro Asp His 1160 Gly Ala Pro Ala Arg Gly Ser Pro His Ser Glu Gly Gly Lys Arg Ser 1175 1180 Pro Glu Pro Asn Lys Thr Ser Val Leu Gly Gly Glu Asp Gly Ile 1190 1195 Glu Pro Val Ser Pro Pro Glu Gly Met Thr Glu Pro Gly His Ser Arg 1210 1205

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			cgg acc cag Arg Thr Gln 5		
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Lys Asn Phe Gly Leu Ile Ala Ser Phe Leu Glu Arg Lys Thr Val Ala 455 Glu Cys Val Leu Tyr Tyr Leu Thr Lys Lys Asn Glu Asn Tyr Lys 475 470 Ser Leu Val Arg Arg Ser Tyr Arg Arg Gly Lys Ser Gln Gln 490 485 505 Pro Arg Ser Ser Gln Glu Glu Lys Asp Glu Lys Glu Lys Glu Lys Glu 520 Ala Glu Lys Glu Glu Glu Lys Pro Glu Val Glu Asn Asp Lys Glu Asp 535 Leu Leu Lys Glu Lys Thr Asp Asp Thr Ser Gly Glu Asp Asn Asp Glu 550 555 Lys Glu Ala Val Ala Ser Lys Gly Arg Lys Thr Ala Asn Ser Gln Gly 570 Arg Arg Lys Gly Arg Ile Thr Arg Ser Met Ala Asn Glu Ala Asn Ser 585 Glu Glu Ala Ile Thr Pro Gln Gln Ser Ala Glu Leu Ala Ser Met Glu 600 Leu Asn Glu Ser Ser Arg Trp Thr Glu Glu Glu Met Glu Thr Ala Lys 615 Lys Gly Leu Leu Glu His Gly Arg Asn Trp Ser Ala Ile Ala Arg Met 635 630 Val Gly Ser Lys Thr Val Ser Gln Cys Lys Asn Phe Tyr Phe Asn Tyr 645 650 Lys Lys Arg Gln Asn Leu Asp Glu Ile Leu Gln Gln His Lys Leu Lys 660 665 Met Glu Lys Glu Arg Asn Ala Arg Arg Lys Lys Lys Ala Pro Ala 680 Ala Ala Ser Glu Glu Ala Ala Phe Pro Pro Val Val Glu Asp Glu Glu 695 700 Met Glu Ala Ser Gly Val Ser Gly Asn Glu Glu Met Val Glu Glu 710 715 Ala Glu Ala Leu His Ala Ser Gly Asn Glu Val Pro Arg Gly Glu Cys 725 730 Ser Gly Pro Ala Thr Val Asn Asn Ser Ser Asp Thr Glu Ser Ile Pro 745 Ser Pro His Thr Glu Ala Ala Lys Asp Thr Gly Gln Asn Gly Pro Lys 760 Pro Pro Ala Thr Leu Gly Ala Asp Gly Pro Pro Pro Gly Pro Pro Thr 775 780 Pro Pro Arg Arg Thr Ser Arg Ala Pro Ile Glu Pro Thr Pro Ala Ser 790 795 Glu Ala Thr Gly Ala Pro Thr Pro Pro Pro Ala Pro Pro Ser Pro Ser 810 Ala Pro Pro Pro Val Val Pro Lys Glu Glu Lys Glu Glu Glu Thr Ala 825 Ala Ala Pro Pro Val Glu Glu Glu Glu Gln Lys Pro Pro Ala Ala 840 Glu Glu Leu Ala Val Asp Thr Gly Lys Ala Glu Glu Pro Val Lys Ser 855 Glu Cys Thr Glu Glu Ala Glu Glu Gly Pro Ala Lys Gly Lys Asp Ala 875 870 Glu Ala Ala Glu Ala Thr Ala Glu Gly Ala Leu Lys Ala Glu Lys Lys 890 Glu Gly Gly Ser Gly Arg Ala Thr Thr Ala Lys Ser Ser Gly Ala Pro 900 905 Gln Asp Ser Asp Ser Ser Ala Thr Cys Ser Ala Asp Glu Val Asp Glu 920

Ala Glu Gly Gly Asp Lys Asn Arg Leu Leu Ser Pro Arg Pro Ser Leu Leu Thr Pro Thr Gly Asp Pro Arg Ala Asn Ala Ser Pro Gln Lys Pro 950 Leu Asp Leu Lys Gln Leu Lys Gln Arg Ala Ala Ile Pro Pro Ile 970 965 Gln Val Thr Lys Val His Glu Pro Pro Arg Glu Asp Ala Ala Pro Thr 985 980 Lys Pro Ala Pro Pro Ala Pro Pro Pro Gln Asn Leu Gln Pro Glu 1000 Ser Asp Ala Pro Gln Gln Pro Gly Ser Ser Pro Arg Gly Lys Ser Arg 1020 1015 Ser Pro Ala Pro Pro Ala Asp Lys Glu Ala Phe Ala Ala Glu Ala Gln 1030 1035 Lys Leu Pro Gly Asp Pro Pro Cys Trp Thr Ser Gly Leu Pro Phe Pro 1045 1050 Val Pro Pro Arg Glu Val Ile Lys Ala Ser Pro His Ala Pro Asp Pro 1065 1060 Ser Ala Phe Ser Tyr Ala Pro Pro Gly His Pro Leu Pro Leu Gly Leu 1075 1080 1085 His Asp Thr Ala Arg Pro Val Leu Pro Arg Pro Pro Thr Ile Ser Asn 1095 1100 Pro Pro Pro Leu Ile Ser Ser Ala Lys His Pro Ser Val Leu Glu Arg 1110 1115 Gln Ile Gly Ala Ile Ser Gln Gly Met Ser Val Gln Leu His Val Pro 1125 1130 Tyr Ser Glu His Ala Lys Ala Pro Val Gly Pro Val Thr Met Gly Leu 1140 1145 1150 Pro Leu Pro Met Asp Pro Lys Lys Leu Ala Pro Phe Ser Gly Val Lys 1160 1165 Gln Glu Gln Leu Ser Pro Arg Gly Gln Ala Gly Pro Pro Glu Ser Leu 1175 1180 Gly Val Pro Thr Ala Gln Glu Ala Ser Val Leu Arg Gly Thr Ala Leu 1190 1195 Gly Ser Val Pro Gly Gly Ser Ile Thr Lys Gly Ile Pro Ser Thr Arg 1205 1210 Val Pro Ser Asp Ser Ala Ile Thr Tyr Arg Gly Ser Ile Thr His Gly 1225 Thr Pro Ala Asp Val Leu Tyr Lys Gly Thr Ile Thr Arg Ile Ile Gly 1240 Glu Asp Ser Pro Ser Arg Leu Asp Arg Gly Arg Glu Asp Ser Leu Pro 1260 1255 Lys Gly His Val Ile Tyr Glu Gly Lys Lys Gly His Val Leu Ser Tyr 1270 1275 Glu Gly Gly Met Ser Val Thr Gln Cys Ser Lys Glu Asp Gly Arg Ser 1285 1290 Ser Ser Gly Pro Pro His Glu Thr Ala Ala Pro Lys Arg Thr Tyr Asp 1305 1300 Met Met Glu Gly Arg Val Gly Arg Ala Ile Ser Ser Ala Ser Ile Glu 1320 1325 Gly Leu Met Gly Arg Ala Ile Pro Pro Glu Arg His Ser Pro His His 1335 1340 Leu Lys Glu Gln His His Ile Arg Gly Ser Ile Thr Gln Gly Ile Pro 1350 1355 Arg Ser Tyr Val Glu Ala Gln Glu Asp Tyr Leu Arg Arg Glu Ala Lys 1365 1370 Leu Leu Lys Arg Glu Gly Thr Pro Pro Pro Pro Pro Pro Ser Arg Asp 1385 Leu Thr Glu Ala Tyr Lys Thr Gln Ala Leu Gly Pro Leu Lys Leu Lys . 1400 1405

Pro Ala His Glu Gly Leu Val Ala Thr Val Lys Glu Ala Gly Arg Ser 1415 Ile His Glu Ile Pro Arg Glu Glu Leu Arg His Thr Pro Glu Leu Pro 1435 1430 Leu Ala Pro Arg Pro Leu Lys Glu Gly Ser Ile Thr Gln Gly Thr Pro 1450 1445 Leu Lys Tyr Asp Thr Gly Ala Ser Thr Thr Gly Ser Lys Lys His Asp 1460 1465 Val Arg Ser Leu Ile Gly Ser Pro Gly Arg Thr Phe Pro Pro Val His 1480 1485 Pro Leu Asp Val Met Ala Asp Ala Arg Ala Leu Glu Arg Ala Cys Tyr 1495 1500 Glu Glu Ser Leu Lys Ser Arg Pro Gly Thr Ala Ser Ser Ser Gly Gly 1510 1515 Ser Ile Ala Arg Gly Ala Pro Val Ile Val Pro Glu Leu Gly Lys Pro 1525 1530 Arg Gln Ser Pro Leu Thr Tyr Glu Asp His Gly Ala Pro Phe Ala Gly 1540 1545 His Leu Pro Arg Gly Ser Pro Val Thr Met Arg Glu Pro Thr Pro Arg 1555 1560 1565 Leu Gln Glu Gly Ser Leu Ser Ser Ser Lys Ala Ser Gln Asp Arg Lys 1575 1580 Leu Thr Ser Thr Pro Arg Glu Ile Ala Lys Ser Pro His Ser Thr Val 1590 1595 Pro Glu His His Pro His Pro Ile Ser Pro Tyr Glu His Leu Leu Arg 1605 1610 Gly Val Ser Gly Val Asp Leu Tyr Arg Ser His Ile Pro Leu Ala Phe 1620 1625 Asp Pro Thr Ser Ile Pro Arg Gly Ile Pro Leu Asp Ala Ala Ala Ala 1635 1640 1645 Tyr Tyr Leu Pro Arg His Leu Ala Pro Asn Pro Thr Tyr Pro His Leu 1650 1655 1660 Tyr Pro Pro Tyr Leu Ile Arg Gly Tyr Pro Asp Thr Ala Ala Leu Glu 1670 1675 Asn Arg Gln Thr Ile Ile Asn Asp Tyr Ile Thr Ser Gln Gln Met His 1685 1690 His Asn Thr Ala Thr Ala Met Ala Gln Arg Ala Asp Met Leu Arg Gly 1705 Leu Ser Pro Arg Glu Ser Ser Leu Ala Leu Asn Tyr Ala Ala Gly Pro 1720 Arg Gly Ile Ile Asp Leu Ser Gln Val Pro His Leu Pro Val Leu Val 1735 Pro Pro Thr Pro Gly Thr Pro Ala Thr Ala Met Asp Arg Leu Ala Tyr 1750 1755 Leu Pro Thr Ala Pro Gln Pro Phe Ser Ser Arg His Ser Ser Ser Pro 1765 1770 Leu Ser Pro Gly Gly Pro Thr His Leu Thr Lys Pro Thr Thr Ser 1785 1780 Ser Ser Glu Arg Glu Arg Asp Arg Asp Arg Glu Arg Asp Arg Asp Arg 1800 Glu Arg Glu Lys Ser Ile Leu Thr Ser Thr Thr Thr Val Glu His Ala 1815 1820 Pro Ile Trp Arg Pro Gly Thr Glu Gln Ser Ser Gly Ser Ser Gly Ser 1835 1830 Ser Gly Gly Gly Gly Ser Ser Ser Arg Pro Ala Ser His Ser His 1845 1850 Ala His Gln His Ser Pro Ile Ser Pro Arg Thr Gln Asp Ala Leu Gln 1865 1870 Gln Arg Pro Ser Val Leu His Asn Thr Gly Met Lys Gly Ile Ile Thr 1880

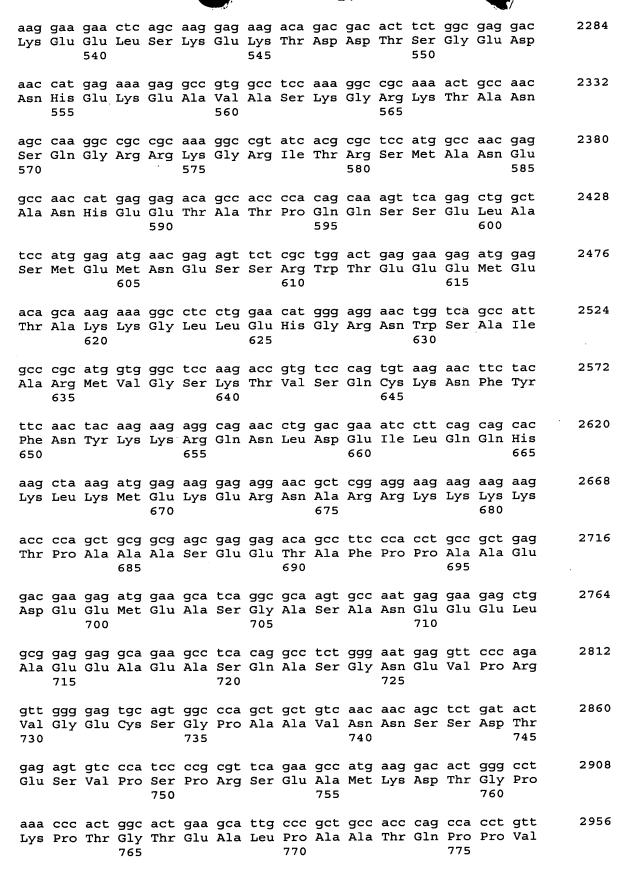
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Asn Pro Leu Asn Ala Ser Ala Ser Leu Pro Ala Ala Met Pro Ile Thr 2375 Ala Ala Asp Gly Arg Ser Asp His Thr Leu Thr Ser Pro Gly Gly Gly 2395 2390 Gly Lys Ala Lys Val Ser Gly Arg Pro Ser Ser Arg Lys Ala Lys Ser 2410 2405 Pro Ala Pro Gly Leu Ala Ser Gly Asp Arg Pro Pro Ser Val Ser Ser 2425 2420 Val His Ser Glu Gly Asp Cys Asn Arg Arg Thr Pro Leu Thr Asn Arg 2445 2440 Val Trp Glu Asp Arg Pro Ser Ser Ala Gly Ser Thr Pro Phe Pro Tyr 2455 2460 Asn Pro Leu Ile Met Arg Leu Gln Ala Gly Val Met Ala Ser Pro Pro 2470 2475 Pro Pro Gly Leu Pro Ala Gly Ser Gly Pro Leu Ala Gly Pro His His 2485 2490 Ala Trp Asp Glu Glu Pro Lys Pro Leu Leu Cys Ser Gln Tyr Glu Thr 2505 2500 Leu Ser Asp Ser Glu 2515 <210> 6 <211> 8388 <212> DNA <213> Mus musculus <220> <221> CDS <222> (626)...(8047) <221> misc feature <222> (1)...(8388) <223> n = A, T, C or G<400> 6 60 cttaaaaaaa aaacccttac ttgtggttaa aggaaaagaa ataaagactt aggaaaaatg taattttcca gggggtacct acacccaaga catatggttc tcaagaggna ctcagcatat 120 180 cactttgatt ccagagaagc tacaaaagtc attaccaaac tccaggctgg aaagcagtgc tcatactaaa tatttaaaca tttaaagacc tgattaagag acatcaaagg ctttatacca 240 ggggcacacc aacagagaca ggctttttca aggataattt atgtctgccc attgtcttct 300 ggcttaggag acatagaggg aaacatcacc taggaaaacc agtaaccaat gtgtaccatc 360 caggagttat tctatgacaa aaccaaaagt tttgttcttg tgtacttctc tgtgcaccat 420 ctttctatat ctatttagaa aacaaaacaa attttggtaa cacgcttgtg tataaagagc 480 540 aggacagcgg tgtcacagat caacctagaa agtaattatt taacgagtaa atgactcata 600 taggacaagg caagctgtga ctttcaacct gttctgtctc gtgccgaatt cggcacgagc 652 caaaqcctac ctqqacccta ccacc atq tca gga tcc aca cag cct gtg gca Met Ser Gly Ser Thr Gln Pro Val Ala cag aca tgg cgg gct gct gag ccc cgc tac cca ccc cat ggc atc tcc 700 Gln Thr Trp Arg Ala Ala Glu Pro Arg Tyr Pro Pro His Gly Ile Ser 10 15 tac ccg gtg cag ata gcc cgg tcc cac acg gac gtg ggg ctg ctt gag 748 Tyr Pro Val Gln Ile Ala Arg Ser His Thr Asp Val Gly Leu Leu Glu 30 796 tac caa cac cac ccc cgt gac tac acc tca cac ctg tca ccc ggt tcc Tyr Gln His His Pro Arg Asp Tyr Thr Ser His Leu Ser Pro Gly Ser 50

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					aaa Lys 335											1660
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gag Glu	aag Lys	gag Glu	gcc Ala 525	gac Asp	aag Lys	gag Glu	gaa Glu	gag Glu 530	aag Lys	cag Gln	gat Asp	gcg Ala	gag Glu 535	aac Asn	gag Glu	2236



cct Pro	cct Pro	cca Pro 780	gaa Glu	gaa Glu	ccg Pro	gca Ala	gta Val 785	gcc Ala	cct Pro	gct Ala	gag Glu	ccc Pro 790	tcc Ser	cca Pro	gtc Val	3004
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													cgg Arg			3100
-			_	_		_			_	_	_	_	agg Arg	_		3148
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_		_	_	_	_	_		_					gaa Glu			3244
													tct Ser			3292
cca Pro 890	ctt Leu	aag Lys	gtg Val	gag Glu	gag Glu 895	gct Ala	ggt Gly	agc Ser	aag Lys	gca Ala 900	gct Ala	gtg Val	acc Thr	aag Lys	ggt Gly 905	3340
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					Pro					Pro			cca Pro		Thr	3628
				Pro					Ser				gga Gly 101	Gly		3676

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ccc act cac cta gct aaa cca act gcc aca tct tca tcg gag cgg gaa Pro Thr His Leu Ala Lys Pro Thr Ala Thr Ser Ser Ser Glu Arg Glu 1755 1760 1765	5932
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aca gtg gag cat gca ccc atc tgg aga cct ggt acg gag cag agc agc Thr Val Glu His Ala Pro Ile Trp Arg Pro Gly Thr Glu Gln Ser Ser 1790 1795 1800	6028
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ctg cac aac acg agc atg aag ggc gtg gtc acc tcc gtg gaa ccc ggc Leu His Asn Thr Ser Met Lys Gly Val Val Thr Ser Val Glu Pro Gly 1835 1840 1845	6172
acg ccc acg gtc ctg agg tgg gcc agg tcc acc tcc acc tct tcg cct Thr Pro Thr Val Leu Arg Trp Ala Arg Ser Thr Ser Thr Ser Ser Pro 1850 1855 1860 1865	6220
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cat gcc agt ccg gac ccg ccg gcg ccc acc tcg gcc tca gat ctg cac His Ala Ser Pro Asp Pro Pro Ala Pro Thr Ser Ala Ser Asp Leu His 1965 1970 1975	6556

cga gaa aag act caa Arg Glu Lys Thr Gli 1980			Leu Glu Leu
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gag ccc atc agc ccc Glu Pro Ile Ser Pro 2010			
ctc tcc aaa cct ctc Leu Ser Lys Pro Leu 203	ı Glu Glu Leu Gl		
ctg cgg cac aag cag Leu Arg His Lys Gli 2045	g cca ggc ccc at n Pro Gly Pro Me 20	t Lys Leu Ser Ala	gag gct gcc 6796 Glu Ala Ala 2055
cat ctc cca cat ctc His Leu Pro His Leu 2060			Ser Ser Ser
cca ctc ctc cag act Pro Leu Leu Gln The 2075	gcc cca ggc at Ala Pro Gly Il 2080	c aaa ggt cac cag e Lys Gly His Gln 2085	agg gtg gtc 6892 Arg Val Val
acc ctg gct cag cac Thr Leu Ala Gln His 2090	c atc agc gag gt s Ile Ser Glu Va 2095	c att acg cag gac l Ile Thr Gln Asp 2100	tac acg cgc 6940 Tyr Thr Arg 2105
cac cac ccg cag cag His His Pro Gln Gln 21:	l Leu Ser Gly Pr		
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gac ctc tac ctc cca Asp Leu Tyr Leu Pro 2140			Arg Gly Ser
ccc cac agt gaa gg Pro His Ser Glu Gl 2155	g ggc aaa agg to y Gly Lys Arg Se 2160	c cca gaa ccc agc r Pro Glu Pro Ser 2165	aaa aca tcg 7132 Lys Thr Ser
gtc ctg ggc agc agc Val Leu Gly Ser Se 2170	c gag gat gcc at c Glu Asp Ala Il 2175	t gag cct gtg tcc e Glu Pro Val Ser 2180	cca cca gag 7180 Pro Pro Glu 2185
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tat cga gac ggg ga Tyr Arg Asp Gly Gl 2205	ı Gln Gly Glu Pr	c agg atg ggt cta o Arg Met Gly Leu 10	gag tct cca 7276 Glu Ser Pro 2215

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aac tcc gcc atg Asn Ser Ala Met 2235		Lys Lys				
aac acc cac aac Asn Thr His Asn 2250				Ile Gly		
acg gaa atc ttc Thr Glu Ile Phe	aac atg ccc Asn Met Pro 2270	gcc atc Ala Ile	act gga Thr Gly 2275	gca ggc Ala Gly	ctt atg Leu Met 2280	Thr
tgt aga agc cag Cys Arg Ser Gln 228	Ala Val Glr	gaa cac Glu His 229	Ala Ser	acc aac Thr Asn	atg ggg Met Gly 2295	cta 7516 Leu
gag gcc att att Glu Ala Ile Ile 2300					Gln Trp	
gag ccc ccg ccg Glu Pro Pro Pro 2315	ctc ggc gcc Leu Gly Ala 232	Asn Ala	ttt aac Phe Asn	cct ctg Pro Leu 2325	aat gcc Asn Ala	agc 7612 Ser
gcc agt ctg ccc Ala Ser Leu Pro 2330				Thr Ala		
agt gac cac gca Ser Asp His Ala						Val
tct ggc aga cct Ser Gly Arg Pro 236	Ser Ser Arg	_	Lys Ser	_		
gcg tcc gga gac Ala Ser Gly Asp 2380	cga ccc cct Arg Pro Pro	tct gtc Ser Val 2385	tcc tca Ser Ser	gta cac Val His 239	Ser Glu	ggg 7804 Gly
gac tgc aat cgc Asp Cys Asn Arg 2395		Leu Thr				
ccc tca tct gca Pro Ser Ser Ala 2410				Asn Pro		
agg cta cag gca Arg Leu Gln Ala						Ala
gca ggc agc ggg Ala Gly Ser Gly 244	Pro Leu Ala	ggt ccc Gly Pro 245	His His	gcc tgg Ala Trp	gat gag Asp Glu 2455	gag 7996 Glu

ccc aag cca ctg ctg tgt tca cag tat gag aca ctc tcg gac agc gag Pro Lys Pro Leu Leu Cys Ser Gln Tyr Glu Thr Leu Ser Asp Ser Glu 2460 2465 2470	8044
tga ccacggattg ggggggagcg gtgccaggtc ccgcacaagg cagaagcagc	8097
*	
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Ser His Thr Asp Val Gly Leu Leu Glu Tyr Gln His His Pro Arg Asp 35 40 45	
Tyr Thr Ser His Leu Ser Pro Gly Ser Ile Ile Gln Pro Gln Arg Arg 50 60	
Arg Pro Ser Leu Leu Ser Glu Phe Gln Pro Gly Ser Glu Arg Ser Gln 65 70 75 80	
Glu Leu His Leu Arg Pro Glu Ser Arg Thr Phe Leu Pro Glu Leu Gly 85 90 95	
Lys Pro Asp Ile Glu Phe Thr Glu Ser Lys Arg Pro Arg Leu Glu Leu 100 105	
Leu Pro Asp Thr Leu Leu Arg Pro Ser Pro Leu Leu Ala Thr Gly Gln 115 120 125	
Pro Ser Gly Ser Glu Asp Leu Thr Lys Asp Arg Ser Leu Ala Gly Lys 130 135 140	
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Glu Leu Ala Pro Ser Arg Leu Ser Lys Glu Glu Leu Ile Gln Asn Met 165 170 175	
Asp Arg Val Asp Arg Glu Ile Thr Met Val Glu Gln Gln Ile Ser Lys 180 185 190	
Leu Lys Lys Gln Gln Gln Leu Glu Glu Glu Ala Ala Lys Pro Pro 195 200 205	
Glu Pro Glu Lys Pro Val Ser Pro Pro Pro Ile Glu Ser Lys His Arg 210 215 220	
Ser Leu Val Gln Ile Ile Tyr Asp Glu Asn Arg Lys Lys Ala Glu Ala	
225 230 235 240	
Ala His Arg Ile Leu Glu Gly Leu Gly Pro Gln Val Glu Leu Pro Leu 245 250 255	
Tyr Asn Gln Pro Ser Asp Thr Arg Gln Tyr His Glu Asn Ile Lys Ile 260 265 . 270	
Asn Gln Ala Met Arg Lys Lys Leu Ile Leu Tyr Phe Lys Arg Arg Asn 275 280 285	
His Ala Arg Lys Gln Trp Glu Gln Arg Phe Cys Gln Arg Tyr Asp Gln 290 295 300	
Leu Met Glu Ala Trp Glu Lys Lys Val Glu Arg Ile Glu Asn Asn Pro	

310 315 305 Arg Arg Arg Ala Lys Glu Ser Lys Val Arg Glu Tyr Tyr Glu Lys Gln 325 330 Phe Pro Glu Ile Arg Lys Gln Arg Glu Leu Gln Glu Arg Met Gln Ser 345 Arg Val Gly Gln Arg Gly Ser Gly Leu Ser Met Ser Ala Ala Arg Ser 360 Glu His Glu Val Ser Glu Ile Ile Asp Gly Leu Ser Glu Gln Glu Asn 375 380 Leu Glu Lys Gln Met Arg Gln Leu Ala Val Ile Pro Pro Met Leu Tyr 390 395 Asp Ala Asp Gln Gln Arg Ile Lys Phe Ile Asn Met Asn Gly Leu Met 405 410 Asp Asp Pro Met Lys Val Tyr Lys Asp Arg Gln Val Thr Asn Met Trp 425 Ser Glu Gln Glu Arg Asp Thr Phe Arg Glu Lys Phe Met Gln His Pro 440 Lys Asn Phe Gly Leu Ile Ala Ser Phe Leu Glu Arg Lys Thr Val Ala 455 Glu Cys Val Leu Tyr Tyr Tyr Leu Thr Lys Lys Asn Glu Asn Tyr Lys 470 475 Ser Leu Val Arg Arg Ser Tyr Arg Arg Gly Lys Ser Gln Gln 485 490 Gln Gln Gln Gln Gln Gln Gln Gln Gln Het Ala Arg Ser Ser 500 505 Gln Glu Glu Lys Glu Lys Glu Lys Glu Lys Glu Ala Asp Lys Glu 520 Glu Glu Lys Gln Asp Ala Glu Asn Glu Lys Glu Glu Leu Ser Lys Glu 535 540 Lys Thr Asp Asp Thr Ser Gly Glu Asp Asn His Glu Lys Glu Ala Val 550 555 Ala Ser Lys Gly Arg Lys Thr Ala Asn Ser Gln Gly Arg Arg Lys Gly 565 570 Arg Ile Thr Arg Ser Met Ala Asn Glu Ala Asn His Glu Glu Thr Ala 585 580 Thr Pro Gln Gln Ser Ser Glu Leu Ala Ser Met Glu Met Asn Glu Ser 600 605 Ser Arg Trp Thr Glu Glu Glu Met Glu Thr Ala Lys Lys Gly Leu Leu 615 620 Glu His Gly Arg Asn Trp Ser Ala Ile Ala Arg Met Val Gly Ser Lys 630 635 Thr Val Ser Gln Cys Lys Asn Phe Tyr Phe Asn Tyr Lys Lys Arg Gln 650 Asn Leu Asp Glu Ile Leu Gln Gln His Lys Leu Lys Met Glu Lys Glu 665 Arg Asn Ala Arg Arg Lys Lys Lys Thr Pro Ala Ala Ala Ser Glu 680 Glu Thr Ala Phe Pro Pro Ala Ala Glu Asp Glu Glu Met Glu Ala Ser 695 700 Gly Ala Ser Ala Asn Glu Glu Glu Leu Ala Glu Glu Ala Glu Ala Ser 710 715 Gln Ala Ser Gly Asn Glu Val Pro Arg Val Gly Glu Cys Ser Gly Pro 730 725 Ala Ala Val Asn Asn Ser Ser Asp Thr Glu Ser Val Pro Ser Pro Arg 745 740 Ser Glu Ala Met Lys Asp Thr Gly Pro Lys Pro Thr Gly Thr Glu Ala 760 765 Leu Pro Ala Ala Thr Gln Pro Pro Val Pro Pro Pro Glu Glu Pro Ala 775 Val Ala Pro Ala Glu Pro Ser Pro Val Pro Asp Ala Ser Gly Pro Pro

795 800 790 785 Ser Pro Glu Pro Ser His His Leu Pro His Pro Arg Leu Leu Trp Thr 805 810 Arg Met Asn Lys Lys Pro Arg Leu Leu Gln Leu Pro Arg Gln Arg Met 820 825 Pro Arg Ser Arg Ser Leu Arg Pro Arg Arg Ser Met Trp Glu Lys Pro 840 Glu Glu Pro Glu Ala Ser Glu Glu Pro Pro Glu Ser Val Lys Ser Asp 855 His Lys Glu Glu Thr Glu Glu Glu Pro Glu Asp Lys Ala Lys Gly Thr 870 875 Glu Ala Ile Glu Thr Val Ser Glu Ala Pro Leu Lys Val Glu Glu Ala 885 890 Gly Ser Lys Ala Ala Val Thr Lys Gly Ser Ser Ser Gly Ala Thr Gln 905 900 Asp Ser Asp Phe Ser Ala Thr Cys Ser Ala Asp Glu Val Asp Glu Pro 925 920 Glu Gly Gly Asp Lys Gly Arg Leu Leu Ser Pro Arg Pro Ser Leu Leu 935 940 Thr Pro Ala Gly Asp Pro Arg Ala Ser Thr Ser Pro Gln Lys Pro Leu 950 955 Asp Leu Lys Gln Leu Lys Gln Arg Ala Ala Ile Pro Pro Ile Gln 970 965 Val Thr Lys Val His Glu Pro Pro Arg Glu Asp Thr Val Pro Pro Lys 980 985 Pro Val Pro Pro Val Pro Pro Pro Thr Gln His Leu Gln Pro Glu Gly 1000 1005 Asp Val Ser Gln Gln Ser Gly Gly Ser Pro Arg Gly Lys Ser Arg Ser 1020 1015 Pro Val Pro Pro Ala Glu Lys Glu Ala Glu Lys Pro Ala Phe Pro 1035 1030 Ala Phe Pro Thr Glu Gly Pro Lys Leu Pro Thr Glu Pro Pro Arg Trp 1045 1050 Ser Ser Gly Leu Pro Phe Pro Ile Pro Pro Arg Glu Val Ile Lys Thr 1070 1060 1065 Ser Pro His Ala Ala Asp Pro Ser Ala Phe Ser Tyr Thr Pro Pro Gly 1080 1085 His Pro Leu Pro Leu Gly Leu His Asp Ser Ala Arg Pro Val Leu Pro 1100 1090 1095 Arg Pro Pro Ile Ser Asn Pro Pro Pro Leu Ile Ser Ser Ala Lys His 1110 1115 Pro Gly Val Leu Glu Arg Gln Leu Gly Ala Ile Ser Gln Gln Gly Met 1125 1130 Ser Val Gln Leu Arg Val Pro His Ser Glu His Ala Lys Ala Pro Met 1145 Gly Pro Leu Thr Met Gly Leu Pro Leu Ala Val Asp Pro Lys Lys Leu 1160 Gly Thr Ala Leu Gly Ser Ala Thr Ser Gly Ser Ile Thr Lys Gly Leu 1175 1180 Pro Ser Thr Arg Ala Ala Asp Gly Pro Ser Tyr Arg Gly Ser Ile Thr 1190 1195 His Gly Thr Pro Ala Asp Val Leu Tyr Lys Gly Thr Ile Ser Arg Ile 1205 1210 Val Gly Glu Asp Ser Pro Ser Arg Leu Asp Arg Ala Arg Glu Asp Thr 1220 1225 Leu Pro Lys Gly His Val Ile Tyr Glu Gly Lys Lys Gly His Val Leu 1240 1245 1235 Ser Tyr Glu Gly Gly Met Ser Val Ser Gln Cys Ser Lys Glu Asp Gly 1260 1255 Arg Ser Ser Ser Gly Pro Pro His Glu Thr Ala Ala Pro Lys Arg Thr

1270 1275 1265 Tyr Asp Met Met Glu Gly Arg Val Gly Arg Thr Val Thr Ser Ala Ser 1285 1290 Ile Glu Gly Leu Met Gly Arg Ala Ile Pro Glu Gln His Ser Pro His 1305 1300 Leu Lys Glu Gln His His Ile Arg Gly Ser Ile Thr Gln Gly Ile Pro 1320 1315 Arg Ser Tyr Val Glu Ala Gln Glu Asp Tyr Leu Arg Arg Glu Ala Lys 1335 1340 Leu Leu Lys Arg Glu Gly Thr Pro Pro Pro Pro Pro Pro Pro Arg Asp 1350 1355 Leu Thr Glu Thr Tyr Lys Pro Arg Pro Leu Asp Pro Leu Gly Pro Leu 1365 1370 Lys Leu Lys Pro Thr His Glu Gly Val Val Ala Thr Val Lys Glu Ala 1380 1385 1390 Gly Arg Ser Ile His Glu Ile Pro Arg Glu Glu Leu Arg Arg Thr Pro 1395 1400 1405 Glu Leu Pro Leu Ala Pro Arg Pro Leu Lys Glu Gly Ser Ile Thr Gln 1410 1415 1420 Gly Thr Pro Leu Lys Tyr Asp Ser Gly Ala Pro Ser Thr Gly Thr Lys 1430 1435 Lys His Asp Val Arg Ser Ile Ile Gly Ser Pro Gly Arg Pro Phe Pro . 1445 1450 Ala Leu His Pro Leu Asp Ile Met Ala Asp Ala Arg Ala Leu Glu Arg 1460 1465 1470 Ala Cys Tyr Glu Glu Ser Leu Lys Ser Arg Ser Gly Thr Ser Ser Gly 1475 1480 1485 Ala Gly Gly Ser Ile Thr Arg Gly Ala Pro Val Val Pro Glu Leu 1500 1490 1495 Gly Lys Pro Arg Gln Ser Pro Leu Thr Tyr Glu Asp His Gly Ala Pro 1510 1515 Phe Thr Ser His Leu Pro Arg Gly Ser Pro Val Thr Thr Arg Glu Pro 1525 1530 Thr Pro Arg Leu Gln Glu Gly Ser Leu Leu Ser Ser Lys Ala Ser Gln 1540 1545 1550 Asp Arg Lys Leu Thr Ser Thr Pro Arg Glu Ile Ala Lys Ser Pro His 1555 1560 1565 Ser Thr Val Pro Glu His His Pro His Pro Ile Ser Pro Tyr Glu His 1575 1580 Leu Leu Arg Gly Val Thr Gly Val Asp Leu Tyr Arg Gly His Ile Pro 1590 1595 Leu Ala Phe Asp Pro Thr Ser Ile Pro Arg Gly Ile Pro Leu Glu Ala 1610 1605 Ala Ala Ala Tyr Tyr Leu Pro Arg His Leu Ala Pro Ser Pro Thr 1625 Tyr Pro His Leu Tyr Pro Pro Tyr Leu Ile Arg Gly Tyr Pro Asp Thr 1640 Ala Ala Leu Glu Asn Arg Gln Thr Ile Ile Asn Asp Tyr Ile Thr Ser 1655 1660 Gln Gln Met His His Asn Ala Ala Ser Ala Met Ala Gln Arg Ala Asp 1670 1675 Met Leu Arg Gly Leu Ser Pro Arg Glu Ser Ser Leu Ala Leu Asn Tyr 1690 1685 Ala Ala Gly Pro Arg Gly Ile Ile Asp Leu Ser Gln Val Pro His Leu 1710 1705 1700 Pro Val Leu Val Pro Pro Thr Pro Gly Thr Pro Ala Thr Ala Ile Asp 1720 1725 Arg Leu Ala Tyr Leu Pro Thr Ala Pro Pro Pro Phe Ser Ser Arg His 1740 1735 Ser Ser Pro Leu Ser Pro Gly Gly Pro Thr His Leu Ala Lys Pro

1745	1750		1755		1760
Thr Ala Thr Ser	Ser Ser Glu 1765		Arg Glu Arg 1770	g Glu Arg	Glu Arg 1775
Asp Lys Ser Ile 178		Thr Thr 1785		His Ala 1790	
Trp Arg Pro Gly 1795	Thr Glu Glr	n Ser Ser 1800	Gly Ala Gly	Gly Ser 1805	Ser Arg
Pro Ala Ser His 1810	Thr His Glr 181		Pro Ile Ser 182		Thr Gln
Asp Ala Leu Gln 1825	Gln Arg Pro	Ser Val	Leu His Ası 1835	Thr Ser	Met Lys 1840
Gly Val Val Thr	Ser Val Glu 1845		Thr Pro Thi 1850	. Val Leu	Arg Trp 1855
Ala Arg Ser Thr 186		Ser Pro 1865		Ala Ala 1870	
Pro Pro Ala Thr 1875	_	1880		1885	
Pro Thr Leu Met 1890	Glu Pro Val		Pro Lys Glu 190		Arg Val
Ala Arg Pro Glu 1905	1910		1915		1920
Lys Pro Pro Gly	1925		1930		1935
Pro Arg Ser Leu 194		Ser Ser 1945		Ala Ile 1950	
Thr Pro Ala Lys 1955	Asn Leu Ala	a Pro His 1960	His Ala Se	Pro Asp 1965	Pro Pro
Ala Pro Thr Ser 1970	197	75	198	30	
Pro Phe Ser Ile 1985	1990		1995		2000
Gly Ala Gly Tyr	2005		2010		2015
Ser Pro Ser Leu 202	:.0	2025	;	203	0
Leu Glu Lys Ser 2035		2040		2045	
Pro Met Lys Leu 2050	Ser Ala Glu 205		His Leu Pro		Arg Pro
Leu Pro Glu Ser 2065	2070		2075		2080
Gly Ile Lys Gly	His Gln Arg 2085		Thr Leu Ala 2090	a Gln His	Ile Ser 2095
Glu Val Ile Thr 210		r Thr Arg 2105		o Gln Gln 211	
Gly Pro Leu Pro 2115	Ala Pro Leu	ı Tyr Ser 2120	Phe Pro Gly	y Ala Ser 2125	Cys Pro
Val Leu Asp Leu 2130	Arg Arg Pro		Asp Leu Ty: 21		Pro Pro
Asp His Gly Thr 2145	2150	-	2155		2160
Arg Ser Pro Glu	Pro Ser Lys 2165	s Thr Ser	Val Leu Gly 2170	y Ser Ser	Glu Asp 2175
Ala Ile Glu Pro 218		o Pro Glu 2185		r Glu Pro 219	
Ala Arg Ser Thr 2195	Ala Tyr Pro	Leu Leu 2200	Tyr Arg As	Gly Glu 2205	Gln Gly
Glu Pro Arg Met 2210	: Gly Leu Glu 22:	ı Ser Pro	Gly Asn Th		Pro Pro
Thr Phe Phe Ser			Asn Ser Ala	a Met Val	Lys Ser

37	
2225 2230 2235 2240	
Lys Lys Gln Glu Ile Asn Lys Lys Leu Asn Thr His Asn Arg Asn Glu 2245 2250 2255	
Pro Glu Tyr Asn Ile Gly Gln Pro Gly Thr Glu Ile Phe Asn Met Pro 2260 2265 2270	
Ala Ile Thr Gly Ala Gly Leu Met Thr Cys Arg Ser Gln Ala Val Gln 2275 2280 2285	
Glu His Ala Ser Thr Asn Met Gly Leu Glu Ala Ile Ile Arg Lys Ala 2290 2295 2300	
Leu Met Gly Lys Tyr Asp Gln Trp Glu Glu Pro Pro Pro Leu Gly Ala 2305 2310 2315 2320	
Asn Ala Phe Asn Pro Leu Asn Ala Ser Ala Ser Leu Pro Ala Ala Ala 2325 2330 2335	
Met Pro Ile Thr Thr Ala Asp Gly Arg Ser Asp His Ala Leu Thr Ser 2340 2345 2350	
Pro Gly Gly Gly Lys Ala Lys Val Ser Gly Arg Pro Ser Ser Arg 2355 2360 2365	
Lys Ala Lys Ser Pro Ala Pro Gly Leu Ala Ser Gly Asp Arg Pro Pro 2370 2375 2380	
Ser Val Ser Ser Val His Ser Glu Gly Asp Cys Asn Arg Arg Thr Pro 2385 2390 2395 2400	
Leu Thr Asn Arg Val Trp Glu Asp Arg Pro Ser Ser Ala Gly Ser Thr 2405 2410 2415	
Pro Phe Pro Tyr Asn Pro Leu Ile Met Arg Leu Gln Ala Gly Val Met 2420 2425 2430	
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Met Ser Gly Ser Thr Gln Pro Val Ala Gln Thr Trp Arg Ala Ala	
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Glu Pro Arg Tyr Pro Pro His Gly Ile Ser Tyr Pro Val Gln Ile Ala	
20 25 30	

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					ata Ile											551
					aac Asn											599
		_		_	cag Gln 85											647
					ccg Pro											695
					cag Gln											743
					agc Ser											791
					agt Ser											839
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					cct Pro											1079
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											tca Ser					1511
											gaa Glu 395					1559
											tgg Trp					1607
											aag Lys					1655
											ctt Leu					1703
											aag Lys					1751
											cct Pro 475					1799
gaa Glu 480	gag Glu	atg Met	gaa Glu	gca Ala	tca Ser 485	ggc Gly	gca Ala	agt Ser	gcc Ala	aat Asn 490	gag Glu	gaa Glu	gag Glu	ctg Leu	gcg Ala 495	1847
gag Glu	gag Glu	gca Ala	gaa Glu	gcc Ala 500	tca Ser	cag Gln	gcc Ala	tct Ser	505 939	Asn	gag Glu	gtt Val	ccc Pro	aga Arg 510	gtt Val	1895

									_	1 0							
	gag Glu															:	1943
	gtc Val															,	1991
	act Thr 545															:	2039
	cca Pro															!	2087
	gcc Ala															:	2135
	cgg Arg																2183
	ccc Pro	_	-		_			_									2231
tcg Ser	atg Met 625	tgg Trp	gaa Glu	aag Lys	cca Pro	gag Glu 630	gag Glu	ccc Pro	gag Glu	gct Ala	tct Ser 635	gag Glu	aag Lys	ccc Pro	ccg Pro		2279
	agt Ser																2327
	aaa Lys																2375
	aag Lys																2423
agc Ser	tca Ser	ggt Gly 690	gcc Ala	acc Thr	cag Gln	gac Asp	agt Ser 695	gac Asp	tcc Ser	agt Ser	gcc Ala	acc Thr 700	tgc Cys	agt Ser	gcc Ala		2471
	gag Glu 705																2519
	agg Arg																2567
tcg Ser	ccc Pro	cag Gln	aag Lys	ccg Pro 740	ctg Leu	gac Asp	ctg Leu	aag Lys	cag Gln 745	ctg Leu	aag Lys	cag Gln	cga Arg	gca Ala 750	gcc Ala		2615

	cct atc gtc Pro Ile Val 755		al His Glu		Glu Asp	2663
	cca aag cca Pro Lys Pro					2711
	gag ggt gac Glu Gly Asp					2759
	cgc agc cca Arg Ser Pro 805					2807
	ttt ccg gct Phe Pro Ala 820					2855
gag ccc cca Glu Pro Pro	cgc tgg tca Arg Trp Ser 835	tcg ggc ct Ser Gly Le 84	eu Pro Phe	ccc atc cct Pro Ile Pro 845	Pro Arg	2903
	aag act tcc Lys Thr Ser					2951
	ccc ggt cac Pro Gly His					2999
	ctg cca cgt Leu Pro Arg 885	Pro Pro Il				3047
	aag cat ccc Lys His Pro 900					3095
tcc cag cag Ser Gln Gln	ggg atg tca Gly Met Ser 915	gtc cag ct Val Gln Le 92	eu Arg Val	cct cac tca Pro His Ser 925	Glu His	3143
gcc aag gcc Ala Lys Ala 930	ccc atg ggc Pro Met Gly	cct ctc ac Pro Leu Th 935	cc atg ggg nr Met Gly	ctg ccc ctt Leu Pro Leu 940	gcc gtg Ala Val	3191
gac cct aag Asp Pro Lys 945	aag ctg ggg Lys Leu Gly	aca gca ct Thr Ala Le 950	ng ggc tcc eu Gly Ser	gcc acc agt Ala Thr Sei 955	gga agc Gly Ser	3239
	ggc ctc ccc Gly Leu Pro 965	Ser Thr Ar				3287
aga ggc tct Arg Gly Ser	atc acc cac Ile Thr His 980	ggc acg co Gly Thr Pr	cc gca gac ro Ala Asp 985	gtc ctc tac Val Leu Tyr	aag ggt Lys Gly 990	3335

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gca Ala	cga Arg	gag Glu 1010	Asp	acc Thr	ctg Leu	ccc Pro	aag Lys 1015	Gly	cat His	gtc Val	atc Ile	tat Tyr 1020	Glu	ggc Gly	aag Lys	3431
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	aag Lys 0					Ser					Pro					3527
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	acc Thr			Ser					Met					Pro		3623
	cac His		Pro					Gln					Gly			3671
	caa Gln 1105	Gly					Tyr					Glu				3719
	cgg Arg 0					Leu					Thr					3767
	cca Pro				Leu					Lys					Asp	3815
	ctg Leu			Leu					Thr					Val		3863
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	cgc Arg 118	Arg					Pro					Pro				3959
	tcc Ser 0					Thr					Asp					4007
	act Thr				Lys					Ser					Pro	4055

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gcc ccc agc ccc acc tac cca cac ctg tac cca cct tac ctc Ala Pro Ser Pro Thr Tyr Pro His Leu Tyr Pro Pro Tyr Let 1410 1415 1420	
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gac tac atc acc tcg cag cag atg cac cac aac gct gcc tc Asp Tyr Ile Thr Ser Gln Gln Met His His Asn Ala Ala Se 1440 1445 1450	
gcc cag cgt gct gac atg ctg agg ggt ctg tca ccg cga ga Ala Gln Arg Ala Asp Met Leu Arg Gly Leu Ser Pro Arg Gl 1460 1465	

									-	-						
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	acc Thr 1505	Ala					Ala					Ala				4919
	agc Ser					Ser					Pro					4967
	cta Leu				Thr					Ser					Glu	5015
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	ggc Gly 1585	Ser					Ser					His				5159
	ccc Pro					Ala					Pro					5207
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	gct Ala		Thr					Thr					Gly			5351
ctt Leu	gaa Glu 1665	Gly	gtc Val	tac Tyr	cct Pro	acc Thr 1670	Leu	atg Met	gag Glu	ccc Pro	gtc Val 1675	Leu	tta Leu	ccc Pro	aag Lys	5399
	acc Thr					Arg					Arg					5447
cat His	gcc Ala	ttt Phe	ctt Leu	acc Thr 1700	Lys	ccc Pro	ccg Pro	ggc Gly	cgg Arg 170	Glu	ccc Pro	gcc Ala	tcc Ser	tca Ser 171	Pro	5495

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aca gcc atc gcc cgc acc cca gca aag aac ctt gca ccc cac cat gcc Thr Ala Ile Ala Arg Thr Pro Ala Lys Asn Leu Ala Pro His His Ala 1730 1735 1740	5591
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ctg ggt tac cac agt gga gct ggc tac agc ccc gat ggg gtg gag ccc Leu Gly Tyr His Ser Gly Ala Gly Tyr Ser Pro Asp Gly Val Glu Pro 1780 1785 1790	5735
atc agc ccg gtg agc tcc ccc agc ctg acc cac gac aag ggg ctc tcc Ile Ser Pro Val Ser Ser Pro Ser Leu Thr His Asp Lys Gly Leu Ser 1795 1800 1805	5783
aaa cct ctg gaa gag cta gag aag agc cac ttg gaa ggg gag ctg cgg Lys Pro Leu Glu Glu Leu Glu Lys Ser His Leu Glu Gly Glu Leu Arg 1810 1815 1820	5831
cac aag cag cca ggc ccc atg aag ctc agc gcg gag gct gcc cat ctc His Lys Gln Pro Gly Pro Met Lys Leu Ser Ala Glu Ala Ala His Leu 1825 1830 1835	5879
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			Gly					Thr			cca Pro		Leu				6311
		Glu					Arg				gag Glu 1995	Ser					6359
	Ser					Phe					act Thr						6407
					Lys					Asn	aag Lys				Thr		6455
				Glu					Ile		cag Gln			Thr			6503
			Met					Gly			ctt Leu		Thr				6551
		Ala					Āla				atg Met 2075	Gly					6599
	Ile					Met					cag Gln)						6647
					Asn					Leu	aat Asn				Ser		6695
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		Ser					Lys				cca Pro 2155	Gly					6839
	Asp					Val					tca Ser						6887
					Leu					Trp	gag Glu				Ser		6935

47	
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agc ggg ccc cta gct ggt ccc cac cac gcc tgg gat gag gag ccc aag Ser Gly Pro Leu Ala Gly Pro His His Ala Trp Asp Glu Glu Pro Lys 2225 2230 2235	7079
cca ctg ctg tgt tca cag tat gag aca ctc tcg gac agc gag tga Pro Leu Leu Cys Ser Gln Tyr Glu Thr Leu Ser Asp Ser Glu * 2240 2245 2250	7124
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Phe Met Gln His Pro Lys Asn Phe Gly Leu Ile Ala Ser Phe Leu Glu

Arg Lys Thr Val Ala Glu Cys Val Leu Tyr Tyr Leu Thr Lys Lys

245 250 Asn Glu Asn Tyr Lys Ser Leu Val Arg Arg Ser Tyr Arg Arg Gly 265 280 Met Ala Arg Ser Ser Gln Glu Glu Lys Glu Glu Lys Glu Lys 295 300 Glu Ala Asp Lys Glu Glu Glu Lys Gln Asp Ala Glu Asn Glu Lys Glu 310 315 Glu Leu Ser Lys Glu Lys Thr Asp Asp Thr Ser Gly Glu Asp Asn Asp 325 330 Glu Lys Glu Ala Val Ala Ser Lys Gly Arg Lys Thr Ala Asn Ser Gln 340 345 Gly Arg Arg Lys Gly Arg Ile Thr Arg Ser Met Ala Asn Glu Ala Asn 360 His Glu Glu Thr Ala Thr Pro Gln Gln Ser Ser Glu Leu Ala Ser Met 375 Glu Met Asn Glu Ser Ser Arg Trp Thr Glu Glu Glu Met Glu Thr Ala 390 395 Lys Lys Gly Leu Leu Glu His Gly Arg Asn Trp Ser Ala Ile Ala Arg 405 410 Met Val Gly Ser Lys Thr Val Ser Gln Cys Lys Asn Phe Tyr Phe Asn 420 425 Tyr Lys Lys Arg Gln Asn Leu Asp Glu Ile Leu Gln Gln His Lys Leu 440 Lys Met Glu Lys Glu Arg Asn Ala Arg Arg Lys Lys Lys Thr Pro 455 460 Ala Ala Ala Ser Glu Glu Thr Ala Phe Pro Pro Ala Ala Glu Asp Glu 470 475 Glu Met Glu Ala Ser Gly Ala Ser Ala Asn Glu Glu Leu Ala Glu 485 490 Glu Ala Glu Ala Ser Gln Ala Ser Gly Asn Glu Val Pro Arg Val Gly 505 500 Glu Cys Ser Gly Pro Ala Ala Val Asn Asn Ser Ser Asp Thr Glu Ser 520 525 Val Pro Ser Pro Arg Ser Glu Ala Thr Lys Asp Thr Gly Pro Lys Pro 535 540 Thr Gly Thr Glu Ala Leu Pro Ala Ala Thr Gln Pro Pro Val Pro Pro 550 555 Pro Glu Glu Pro Ala Val Ala Pro Ala Glu Pro Ser Pro Val Pro Asp 565 570 Ala Ser Gly Pro Pro Ser Pro Glu Pro Ser His His Leu Pro His Pro 580 585 Arg Leu Leu Trp Thr Arg Met Asn Lys Lys Pro Arg Leu Leu Gln Leu 600 Pro Arg Gln Arg Met Pro Arg Ser Arg Ser Leu Arg Pro Arg Arg Ser 615 620 Met Trp Glu Lys Pro Glu Glu Pro Glu Ala Ser Glu Lys Pro Pro Lys 630 635 Ser Val Lys Ser Asp His Lys Lys Glu Thr Glu Glu Glu Pro Glu Asp 645 650 Lys Ala Lys Gly Thr Glu Ala Ile Glu Thr Val Ser Glu Ala Pro Leu 665 Lys Val Glu Lys Ala Gly Ser Lys Ala Ala Val Thr Lys Gly Ser Ser 680 Ser Gly Ala Thr Gln Asp Ser Asp Ser Ser Ala Thr Cys Ser Ala Asp 695 700 Glu Val Asp Glu Pro Glu Gly Gly Asp Lys Gly Arg Leu Leu Ser Pro 710 715 Arg Pro Ser Leu Leu Thr Pro Ala Gly Asp Pro Arg Ala Ser Thr Ser

Pro Gln Lys Pro Leu Asp Leu Lys Gln Leu Lys Gln Arg Ala Ala. Ile Pro Pro Ile Val Thr Lys Val His Glu Pro Pro Arg Glu Asp Thr Val Pro Pro Lys Pro Val Pro Pro Val Pro Pro Pro Thr Gln His Leu Gln Pro Glu Gly Asp Val Ser Gln Gln Ser Gly Gly Ser Pro Arg Gly Lys Ser Arg Ser Pro Val Pro Pro Ala Glu Lys Glu Ala Glu Lys Pro Ala Phe Phe Pro Ala Phe Pro Thr Glu Gly Pro Lys Leu Pro Thr Glu Pro Pro Arg Trp Ser Ser Gly Leu Pro Phe Pro Ile Pro Pro Arg Glu Val Ile Lys Thr Ser Pro His Ala Ala Asp Pro Ser Ala Phe Ser Tyr Thr Pro Pro Gly His Pro Leu Pro Leu Gly Leu His Asp Ser Ala Arg Pro Val Leu Pro Arg Pro Pro Ile Ser Asn Pro Pro Pro Leu Ile Ser Ser Ala Lys His Pro Gly Val Leu Glu Arg Gln Leu Gly Ala Ile Ser Gln Gln Gly Met Ser Val Gln Leu Arg Val Pro His Ser Glu His Ala Lys Ala Pro Met Gly Pro Leu Thr Met Gly Leu Pro Leu Ala Val Asp Pro Lys Lys Leu Gly Thr Ala Leu Gly Ser Ala Thr Ser Gly Ser Ile Thr Lys Gly Leu Pro Ser Thr Arg Ala Ala Asp Gly Pro Ser Tyr Arg Gly Ser Ile Thr His Gly Thr Pro Ala Asp Val Leu Tyr Lys Gly Thr Ile Ser Arg Ile Val Gly Glu Asp Ser Pro Ser Arg Leu Asp Arg Ala Arg Glu Asp Thr Leu Pro Lys Gly His Val Ile Tyr Glu Gly Lys Lys Gly His Val Leu Ser Tyr Glu Gly Gly Met Ser Val Ser Gln Cys Ser Lys Glu Asp Gly Arg Ser Ser Ser Gly Pro Pro His Glu Thr Ala Ala Pro Lys Arg Thr Tyr Asp Met Met Glu Gly Arg Val Gly Arg Thr Val Thr Ser Ala Ser Ile Glu Gly Leu Met Gly Arg Ala Ile Pro Glu Gln His Ser Pro His Leu Lys Glu Gln His His Ile Arg Gly Ser Ile Thr Gln Gly Ile Pro Arg Ser Tyr Val Glu Ala Gln Glu Asp Tyr Leu Arg Arg Glu Ala Lys Leu Leu Lys Arg Glu Gly Thr Pro Pro Pro Pro Pro Pro Arg Asp Leu Thr Glu Thr Tyr Lys Pro Arg Pro Leu Asp Pro Leu Gly Pro Leu Lys Leu Lys Pro Thr His Glu Gly Val Val Ala Thr Val Lys Glu Ala Gly Arg Ser Ile His Glu Ile Pro Arg Glu Glu Leu Arg Arg Thr Pro Glu Leu Pro Leu Ala Pro Arg Pro Leu Lys Glu Gly Ser Ile Thr Gln Gly Thr Pro Leu Lys Tyr Asp Ser Gly Ala Pro Ser

1205 1210 Thr Gly Thr Lys Lys His Asp Val Arg Ser Ile Ile Gly Ser Pro Gly 1220 1225 Arg Pro Phe Pro Ala Leu His Pro Leu Asp Ile Met Ala Asp Ala Arg 1240 Ala Leu Glu Arg Ala Cys Tyr Glu Glu Ser Leu Lys Ser Arg Ser Gly 1255 1260 Thr Ser Ser Gly Ala Gly Gly Ser Ile Thr Arg Gly Ala Pro Val Val 1275 1270 Val Pro Glu Leu Gly Lys Pro Arg Gln Ser Pro Leu Thr Tyr Glu Asp 1290 1295 1285 His Gly Ala Pro Phe Thr Ser His Leu Pro Arg Gly Ser Pro Val Thr 1305 1300 Thr Arg Glu Pro Thr Pro Arg Leu Gln Glu Gly Ser Leu Leu Ser Ser 1315 1320 Lys Ala Ser Gln Asp Arg Lys Leu Thr Ser Thr Pro Arg Glu Ile Ala 1330 1335 1340 Lys Ser Pro His Ser Thr Val Pro Glu His His Pro His Pro Ile Ser 1350 1355 Pro Tyr Glu His Leu Leu Arg Gly Val Thr Gly Val Asp Leu Tyr Arg 1365 1370 Gly His Ile Pro Leu Ala Phe Asp Pro Thr Ser Ile Pro Arg Gly Ile 1385 1380 1390 Pro Leu Glu Ala Ala Ala Ala Tyr Tyr Leu Pro Arg His Leu Ala 1395 1400 1405 Pro Ser Pro Thr Tyr Pro His Leu Tyr Pro Pro Tyr Leu Ile Arg Gly 1410 1415 1420 Tyr Pro Asp Thr Ala Ala Leu Glu Asn Arg Gln Thr Ile Ile Asn Asp 1430 1435 Tyr Ile Thr Ser Gln Gln Met His His Asn Ala Ala Ser Ala Met Ala 1445 1450 1455 Gln Arg Ala Asp Met Leu Arg Gly Leu Ser Pro Arg Glu Ser Ser Leu 1465 1460 Ala Leu Asn Tyr Ala Ala Gly Pro Arg Gly Ile Ile Asp Leu Ser Gln 1475 1480 1485 Val Pro His Leu Pro Val Leu Val Pro Pro Thr Pro Gly Thr Pro Ala 1490 1495 1500 Thr Ala Ile Asp Arg Leu Ala Tyr Leu Pro Thr Ala Pro Pro Phe 1510 1515 Ser Ser Arg His Ser Ser Ser Pro Leu Ser Pro Gly Gly Pro Thr His 1525 1530 Leu Ala Lys Pro Thr Ala Thr Ser Ser Ser Glu Arg Glu Arg Glu Arg 1540 1545 Glu Arg Glu Arg Asp Lys Ser Ile Leu Thr Ser Thr Thr Val Glu 1555 1560 1565 His Ala Pro Ile Trp Arg Pro Gly Thr Glu Gln Ser Ser Gly Ala Gly 1575 1580 Gly Ser Ser Arg Pro Ala Ser His Thr His Gln His Ser Pro Ile Ser 1590 1595 Pro Arg Thr Gln Asp Ala Leu Gln Gln Arg Pro Ser Val Leu His Asn 1605 1610 Thr Ser Met Lys Gly Val Val Thr Ser Val Glu Pro Gly Thr Pro Thr 1625 Val Leu Arg Trp Ala Arg Ser Thr Ser Thr Ser Pro Val Arg Pro 1640 1645 Ala Ala Thr Phe Pro Pro Ala Thr His Cys Pro Leu Gly Gly Thr Leu 1655 1660 Glu Gly Val Tyr Pro Thr Leu Met Glu Pro Val Leu Leu Pro Lys Glu 1675 1670 Thr Ser Arg Val Ala Arg Pro Glu Arg Ala Arg Val Asp Ala Gly His

		51	
	1685	1690	1695
Ala Phe Leu Thr 170		Gly Arg Glu Pro Al 1705	a Ser Ser Pro Ser 1710
		Leu Ala Pro Pro Se 1720	
		Lys Asn Leu Ala Pr	o His His Ala Ser
Pro Asp Pro Pro		Ser Ala Ser Asp Le	u His Arg Glu Lys
1745 Thr Gln Ser Lys		1755 Ile Gln Glu Leu Gl	
		1770 Tyr Ser Pro Asp Gl	=
		1785 Leu Thr His Asp Ly	_
		1800 Ser His Leu Glu Gl	-
1810	181!	5 Leu Ser Ala Glu Al	
1825	1830	1835	1840
His Leu Arg Pro		Ser Gln Pro Ser Se 1850	
Gln Thr Ala Pro		Gly His Gln Arg Va 1865	
		Thr Gln Asp Tyr Th	
Gln Gln Leu Ser 1890	Gly Pro Leu 1899	Pro Ala Pro Leu Ty	
Ala Ser Cys Pro 1905	Val Leu Asp 1910	Leu Arg Arg Pro Pro 1915	Ser Asp Leu Tyr 1920
Leu Pro Pro Pro	Asp His Gly 1925	Thr Pro Ala Arg Gl	y Ser Pro His Ser 1935
Glu Gly Gly Lys		Glu Pro Ser Lys Th 1945	r Ser Val Leu Gly 1950
Ser Ser Glu Asp 1955	Ala Ile Glu	Pro Val Ser Pro Pro 1960	
Glu Pro Gly His 1970	Ala Arg Ser	Thr Ala Tyr Pro Le	
Gly Glu Gln Gly 1985	Glu Pro Arg 1990	Met Gly Leu Glu Se 1995	r Pro Gly Asn Thr 2000
Ser Gln Pro Pro	Thr Phe Phe 2005	Ser Lys Leu Thr Gl	a Ser Asn Ser Ala 2015
Met Val Lys Ser 2020		Glu Ile Asn Lys Lys 2025	s Leu Asn Thr His 2030
Asn Arg Asn Glu 2035	Pro Glu Tyr	Asn Ile Gly Gln Pro 2040	Gly Thr Glu Ile 2045
Phe Asn Met Pro 2050	Ala Ile Thr	Gly Ala Gly Leu Me	
Gln Ala Val Gln	Glu His Ala	Ser Thr Asn Met Gl	y Leu Glu Ala Ile
2065	2070	2075	2080
	2085	Lys Tyr Asp Gln Try 2090	2095
2100	0	Asn Pro Leu Asn Ala 2105	2110
Pro Ala Ala Ala 2115	Met Pro Ile	Thr Thr Ala Asp Gly 2120	y Arg Ser Asp His 2125
Ala Leu Thr Ser 2130	Pro Gly Gly 2135	Gly Gly Lys Ala Lys 5 214	
Pro Ser Ser Arg 2145	Lys Ala Lys 2150	Ser Pro Ala Pro Gly 2155	y Leu Ala Ser Gly 2160
Asp Arg Pro Pro	Ser Val Ser	Ser Val His Ser Gl	Gly Asp Cys Asn

2165 2170 2175										
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Ala Gly Ser Thr Pro Phe Pro Tyr Asn Pro Leu Ile Met Arg Leu Gln 2195 2200 2205										
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Met Ser Ser Gly Tyr Pro Pro Asn Gln Gly Ala Phe Ser Thr Glu 1 5 10 15										
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cgc cac cag cag gag ttc gca gtc cct gat tat cgt tcc tct cat ctt Arg His Gln Gln Glu Phe Ala Val Pro Asp Tyr Arg Ser Ser His Leu 35 40 45	384									
gaa gtg agt cag gca tca cag ctt ttg cag caa cag cag caa cag Glu Val Ser Gln Ala Ser Gln Leu Leu Gln Gln Gln Gln Gln Gln 50 55 60	432									
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tta gtg cac ccg ctg cca gaa ggg ctg agg gct tct gca gat gct aag Leu Val His Pro Leu Pro Glu Gly Leu Arg Ala Ser Ala Asp Ala Lys	672									

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	gat Asp															720
	Gly aaa															768
	aag Lys															816
	aaa Lys															864
	gaa Glu 210															912
	cct Pro															960
	gag Glu															1008
	ggc Gly			_	_	_		_			_			_		1056
	gtg Val															1104
	att Ile 290															1152
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	gtg Val		_		-							_		_	_	1248
	aca Thr		_			_	_	_			_		_			1296
	gaa Glu															1344
	tca Ser															1392

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					cag Gln 390										1440
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					aat Asn						-	_			1584
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	_			_	agt Ser 470	_		_	_	_	_				1680
	_				aat Asn			_		_	_				1728
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					gaa Glu										1824
					gat Asp										1872
					gaa Glu 550										1920
					caa Gln										1968
					aag Lys										2016
					gct Ala										2064
					cca Pro										2112

							atg Met							2160
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							ttc Phe 665							2256
						_	cag Gln			_			_	2304
							tct Ser							2352
		-	_		_	_	gat Asp		_	_		_	_	2400
							gaa Glu							2448
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							gat Asp							2592
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	_				_		gaa Glu 825				_			2736
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					tca Ser										2976
		_			ccg Pro						_	_	_		3024
	_		_	_	gct Ala 935	_				_	_		_		3072
_					gga Gly				_			_			3120
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		Thr			agt Ser		Asn				_	Val		_	3264
	Pro				ata Ile 1015	Thr					Gly				3312
Thr					agg Arg)					Leu					3360
				Ser	gaa Glu				Phe					Ser	3408
			Thr		ggc Gly			Leu					Gln		3456
		Gln	_		ccc Pro	_	Pro		_			Ile			3504
 _				_	gaa Glu		_			_		_			3552

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atc aag cag gaa Ile Lys Gln Glu 1105					u
ggt ctg ttg gtc Gly Leu Leu Val			Val Val Arg		
gga gcc ata caa Gly Ala Ile Gln 114	Glu Gly Ser				
att tca gtg gag Ile Ser Val Glu 1155				Thr Gln Gl	
acc ccg gct ctg Thr Pro Ala Leu 1170		Gly Ile Pro			
ggg tcc att tcg Gly Ser Ile Ser 1185					Y
aga gag gaa gct Arg Glu Glu Ala	_		Ile Tyr Glu		
gga cat atc ttg Gly His Ile Leu 122	Ser Tyr Asp				
agg agt cca aga Arg Ser Pro Arg 1235	_	-		Ser Tyr Gl	
tca gtg gaa gga Ser Val Glu Gly 1250		Gln Gly Met			
gta tca gca ccg Val Ser Ala Pro 1265					- Y
agt cct cat tct Ser Pro His Ser			Val Leu Ser		
atg cag ggg aca Met Gln Gly Thr 1300	Pro Arg Ala				
aaa tat ccc aaa Lys Tyr Pro Lys 1315				Arg Ala Phe	
gaa ggt gcc att Glu Gly Ala Ile					

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ggg cct agc aaa cta tcc cgt gga at Gly Pro Ser Lys Leu Ser Arg Gly Me 1410 1415	
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caa aac acc atg tcc aga ggc tca co Gln Asn Thr Met Ser Arg Gly Ser Pr 1490 1495	
gtt aca att cct cct aac aag tct ac Val Thr Ile Pro Pro Asn Lys Ser Th 1505 1510	
Val Thr Ile Pro Pro Asn Lys Ser Th	r Asn His Glu Arg Lys Ser Thr 1515 1520 cc cca gcg aag tct cca gtg cct 4848
Val Thr Ile Pro Pro Asn Lys Ser The 1505 1510 ctg acc cct acc cag agg gaa agt at Leu Thr Pro Thr Gln Arg Glu Ser Il 1525 ggg gtg gac cct gtc gtg agc cac ag Gly Val Asp Pro Val Val Ser His Se	ar Asn His Glu Arg Lys Ser Thr 1515 1520 C cca gcg aag tct cca gtg cct 4848 Re Pro Ala Lys Ser Pro Val Pro 1530 1535 gt ccg ttt gat ccc cat cac aga 4896
Val Thr Ile Pro Pro Asn Lys Ser The 1505 1510 ctg acc cct acc cag agg gaa agt at Leu Thr Pro Thr Gln Arg Glu Ser Il 1525 ggg gtg gac cct gtc gtg agc cac ag Gly Val Asp Pro Val Val Ser His Se	ar Asn His Glu Arg Lys Ser Thr 1515 1520 15 cc cca gcg aag tct cca gtg cct 1530 1535 154 ccg ttt gat ccc cat cac aga 1550 1550 159 agc cac ctg ccc acg caa ttg 1544

tac ctg ttt cag aga cag ctt tca cca act cca ggt tac cca agt cag Tyr Leu Phe Gln Arg Gln Leu Ser Pro Thr Pro Gly Tyr Pro Ser Gln 1585 1590 1595 1600	5040
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cct cat cca ggg gga aca agc act cct ccc atg gac aga atc act tat Pro His Pro Gly Gly Thr Ser Thr Pro Pro Met Asp Arg Ile Thr Tyr 1665 1670 1675 1680	5280
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gta aga act cag gag acc atg ttg caa cag aga ccc agt gtt ttc caa	
Val Arg Thr Glu Thr Met Leu Gln Gln Arg Pro Ser Val Phe Gln 1765 1770 1775	5568
Val Arg Thr Gln Glu Thr Met Leu Gln Gln Arg Pro Ser Val Phe Gln	5568 5616
Val Arg Thr Gln Glu Thr Met Leu Gln Gln Arg Pro Ser Val Phe Gln 1765 1770 1775 gga acc aat gga acc agt gta atc aca cct ttg gat cca act gct cag Gly Thr Asn Gly Thr Ser Val Ile Thr Pro Leu Asp Pro Thr Ala Gln	

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acc aga ggg aag Thr Arg Gly Lys			Asn Phe Ile		6048
atc acc cgg caa Ile Thr Arg Gln 194	Ile Ala Ser				6096
caa agt tca gac Gln Ser Ser Asp 1955				Tyr Glu Thr	6144
cct agc gat gct Pro Ser Asp Ala 1970		Ile Ser Pro			6192
ccc cag gag aaa Pro Gln Glu Lys 1985					6240
caa gcg gaa aat Gln Ala Glu Asn			Glu Gly Pro		6288
tat cga cca cag Tyr Arg Pro Gln 202	Gln Glu Ser		_	_	6336
tct tca cag gca Ser Ser Gln Ala 2035				His Arg Leu	6384
atc aca ctt gct Ile Thr Leu Ala					6432

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aga aat caa gtt tcc tcg cag act Arg Asn Gln Val Ser Ser Gln Th 2065 2070	Pro Gln Gln Pro Pro Thr Ser	
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gga agc ttt gat gac aaa gtt gag Gly Ser Phe Asp Asp Lys Val Glu 2275 228	Asp His Gly Val Val Met Ser G	
cct atg gga gta gtg cct ggt act Pro Met Gly Val Val Pro Gly Thr		

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2300

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ggt gag aca cga aga gag gaa ggg gac cca tca cct cat tca gga gga 7200 Gly Glu Thr Arg Arg Glu Glu Gly Asp Pro Ser Pro His Ser Gly Gly 2305 2310 gtt tgc aaa cca aag ctg atc agc aag tca aac agc agg aaa tct aag 7248 Val Cys Lys Pro Lys Leu Ile Ser Lys Ser Asn Ser Arg Lys Ser Lys 2325 tet eet ata eet ggg caa gge tae tta gga aeg gaa egg eee tet tea 7296 Ser Pro Ile Pro Gly Gln Gly Tyr Leu Gly Thr Glu Arg Pro Ser Ser gtc tcc tct gta cat tca gaa ggg gat tac cat agg cag acg cca ggg 7344 Val Ser Ser Val His Ser Glu Gly Asp Tyr His Arg Gln Thr Pro Gly 2355 2360 tgg gcc tgg gaa gac agg ccc tct tca aca ggc tca act cag ttt cct 7392 Trp Ala Trp Glu Asp Arg Pro Ser Ser Thr Gly Ser Thr Gln Phe Pro 2375 tat aac cct ctg act atg cgg atg ctc agc agt act cca cca aca ccg 7440 Tyr Asn Pro Leu Thr Met Arg Met Leu Ser Ser Thr Pro Pro Thr Pro 2390 2395 att gca tgt gct ccc tct gcg gtg aac caa gca gct cct cac caa cag 7488 Ile Ala Cys Ala Pro Ser Ala Val Asn Gln Ala Ala Pro His Gln Gln 2405 2410 aac agg atc tgg gag cga gag cct gcc cca ctg ctc tca gca cag tac 7536 Asn Arg Ile Trp Glu Arg Glu Pro Ala Pro Leu Leu Ser Ala Gln Tyr 2420 2425 gag acc ctg tcg gat agt gat gac tga actgcacaaa gtgaggggaa 7583 Glu Thr Leu Ser Asp Ser Asp Asp 2435 2440 cagggtgcag gagagggatc tctagttttt gtggtttaat ttttagtagc aggtcaaaaa 7643 cctgccctcc tgtgacttat tccctgagac ttttcaggag agccagccca cagatgatga 7703 agaaatgatg gaagttcatt tggagagtca aatgggaaaa aaacaaacaa aaaactgcct 7763 ttgatacagg caattcagtg gactataata atagtggagg gttgagatgt agagttttta 7823 aaaagtgaac agttgctgtt cttacatctg taaagaaaac cataatgtct ttaaatcact 7883 cttctgtaaa tagatgacct ttttgcagtg taaaaaaaaa aaaaaaaaa aaaaaaa 7940 <210> 11 <211> 2440 <212> PRT <213> Homo sapiens <400> 11 Met Ser Ser Gly Tyr Pro Pro Asn Gln Gly Ala Phe Ser Thr Glu 10 Gln Ser Arg Tyr Pro Pro His Ser Val Gln Tyr Thr Phe Pro Asn Thr 25 Arg His Gln Gln Glu Phe Ala Val Pro Asp Tyr Arg Ser Ser His Leu 40 Glu Val Ser Gln Ala Ser Gln Leu Leu Gln Gln Gln Gln Gln Gln

Leu Arg Arg Arg Pro Ser Leu Leu Ser Glu Phe His Pro Gly Ser Asp

65 70 75 Arg Pro Gln Glu Arg Arg Thr Ser Tyr Glu Pro Phe His Pro Gly Pro 90 Ser Pro Val Asp His Asp Ser Leu Glu Ser Lys Arg Pro Arg Leu Glu 105 Gln Val Ser Asp Ser His Phe Gln Arg Val Ser Ala Ala Val Leu Pro 120 Leu Val His Pro Leu Pro Glu Gly Leu Arg Ala Ser Ala Asp Ala Lys 135 Lys Asp Pro Ala Phe Gly Gly Lys His Glu Ala Pro Ser Pro Ile 150 155 Ser Gly Gln Pro Cys Gly Asp Asp Gln Asn Ala Ser Pro Ser Lys Leu 165 170 Ser Lys Glu Glu Leu Ile Gln Ser Met Asp Arg Val Asp Arg Glu Ile 185 Ala Lys Val Glu Gln Gln Ile Leu Lys Leu Lys Lys Lys Gln Gln Gln 200 Leu Glu Glu Ala Ala Lys Pro Pro Glu Pro Glu Lys Pro Val Ser 215 Pro Pro Pro Val Glu Gln Lys His Arg Ser Ile Val Gln Ile Ile Tyr 230 235 Asp Glu Asn Arg Lys Lys Ala Glu Glu Ala His Lys Ile Phe Glu Gly 245 250 Leu Gly Pro Lys Val Glu Leu Pro Leu Tyr Asn Gln Pro Ser Asp Thr 265 Lys Val Tyr His Glu Asn Ile Lys Thr Asn Gln Val Met Arg Lys Lys 275 280 Leu Ile Leu Phe Phe Lys Arg Arg Asn His Ala Arg Lys Gln Arg Glu 295 300 Gln Lys Ile Cys Gln Arg Tyr Asp Gln Leu Met Glu Ala Trp Glu Lys 310 315 Lys Val Asp Arg Ile Glu Asn Asn Pro Arg Arg Lys Ala Lys Glu Ser 325 330 Lys Thr Arg Glu Tyr Tyr Glu Lys Gln Phe Pro Glu Ile Arg Lys Gln 345 340 Arg Glu Gln Gln Glu Arg Phe Gln Arg Val Gly Gln Arg Gly Ala Gly 360 365 Leu Ser Ala Thr Ile Ala Arg Ser Glu His Glu Ile Ser Glu Ile Ile 375 380 Asp Gly Leu Ser Glu Gln Glu Asn Asn Glu Lys Gln Met Arg Gln Leu 390 395 Ser Val Ile Pro Pro Met Met Phe Asp Ala Glu Gln Arg Arg Val Lys 405 410 Phe Ile Asn Met Asn Gly Leu Met Glu Asp Pro Met Lys Val Tyr Lys 420 425 Asp Arg Gln Phe Met Asn Val Trp Thr Asp His Glu Lys Glu Ile Phe 440 Lys Asp Lys Phe Ile Gln His Pro Lys Asn Phe Gly Leu Ile Ala Ser 455 460 Tyr Leu Glu Arg Lys Ser Val Pro Asp Cys Val Leu Tyr Tyr Leu 470 475 Thr Lys Lys Asn Glu Asn Tyr Lys Ala Leu Val Arg Arg Asn Tyr Gly 485 490 Lys Arg Arg Gly Arg Asn Gln Gln Ile Ala Arg Pro Ser Gln Glu Glu 505 Lys Val Glu Glu Lys Glu Glu Asp Lys Ala Glu Lys Thr Glu Lys Lys 520 Glu Glu Lys Lys Asp Glu Glu Glu Lys Asp Glu Lys Glu Asp Ser 535 540 Lys Glu Asn Thr Lys Glu Lys Asp Lys Ile Asp Gly Thr Ala Glu Glu

Thr Glu Glu Arg Glu Gln Ala Thr Pro Arg Gly Arg Lys Thr Ala Asn Ser Gln Gly Arg Arg Lys Gly Arg Ile Thr Arg Ser Met Thr Asn Glu Ala Ala Ala Ser Ala Ala Ala Ala Ala Thr Glu Glu Pro Pro Pro Pro Leu Pro Pro Pro Glu Pro Ile Ser Thr Glu Pro Val Glu Thr Ser Arg Trp Thr Glu Glu Glu Met Glu Val Ala Lys Lys Gly Leu Val Glu His Gly Arg Asn Trp Ala Ala Ile Ala Lys Met Val Gly Thr Lys Ser Glu Ala Gln Cys Lys Asn Phe Tyr Phe Asn Tyr Lys Arg Arg His Asn Leu Asp Asn Leu Leu Gln Gln His Lys Gln Lys Thr Ser Arg Lys Pro Arg Glu Glu Arg Asp Val Ser Gln Cys Glu Ser Val Ala Ser Thr Val Ser Ala Gln Glu Asp Glu Asp Ile Glu Ala Ser Asn Glu Glu Glu Asn Pro Glu Asp Ser Glu Val Glu Ala Val Lys Pro Ser Glu Asp Ser Pro Glu Asn Ala Thr Ser Arg Gly Asn Thr Glu Pro Ala Val Glu Leu Glu Pro Thr Thr Glu Thr Ala Pro Ser Thr Ser Pro Ser Leu Ala Val Pro Ser Thr Lys Pro Ala Glu Asp Glu Ser Val Glu Thr Gln Val Asn Asp Ser Ile Ser Ala Glu Thr Ala Glu Gln Met Asp Val Asp Gln Gln Glu His Ser Ala Glu Glu Gly Ser Val Cys Asp Pro Pro Pro Ala Thr Lys Ala Asp Ser Val Asp Val Glu Val Arg Val Pro Glu Asn His Ala Ser Lys Val Glu Gly Asp Asn Thr Lys Glu Arg Asp Leu Asp Arg Ala Ser Glu Lys Val Glu Pro Arg Asp Glu Asp Leu Val Val Ala Gln Gln Ile Asn Ala Gln Arg Pro Glu Pro Gln Ser Asp Asn Asp Ser Ser Ala Thr Cys Ser Ala Asp Glu Asp Val Asp Gly Glu Pro Glu Arg Gln Arg Met Phe Pro Met Asp Ser Lys Pro Ser Leu Leu Asn Pro Thr Gly Ser Ile Leu Val Ser Ser Pro Leu Lys Pro Asn Pro Leu Asp Leu Pro Gln Leu Gln His Arg Ala Ala Val Ile Pro Pro Met Val Ser Cys Thr Pro Cys Asn Ile Pro Ile Gly Thr Pro Val Ser Gly Tyr Ala Leu Tyr Gln Arg His Ile Lys Ala Met His Glu Ser Ala Leu Leu Glu Gln Arg Gln Arg Gln Glu Gln Ile Asp Leu Glu Cys Arg Ser Ser Thr Ser . 985 Pro Cys Gly Thr Ser Lys Ser Pro Asn Arg Glu Trp Glu Val Leu Gln Pro Ala Pro His Gln Leu Ile Thr Asn Leu Pro Glu Gly Val Arg Leu Pro Thr Thr Arg Pro Thr Arg Pro Pro Pro Leu Ile Pro Ser Ser

1025 1030 1035 Lys Thr Thr Val Ala Ser Glu Lys Pro Ser Phe Ile Met Gly Gly Ser 1045 1050 Ile Ser Gln Gly Thr Pro Gly Thr Tyr Leu Thr Ser His Asn Gln Ala 1065 1060 Ser Tyr Thr Gln Glu Thr Pro Lys Pro Ser Val Gly Ser Ile Ser Leu 1080 1085 Gly Leu Pro Arg Gln Gln Glu Ser Ala Lys Ser Ala Thr Leu Pro Tyr 1095 1100 Ile Lys Gln Glu Glu Phe Ser Pro Arg Ser Gln Asn Ser Gln Pro Glu 1110 1115 Gly Leu Leu Val Arg Ala Gln His Glu Gly Val Val Arg Gly Thr Ala 1125 1130 Gly Ala Ile Gln Glu Gly Ser Ile Thr Arg Gly Thr Pro Thr Ser Lys 1145 . 1150 1140 Ile Ser Val Glu Ser Ile Pro Ser Leu Arg Gly Ser Ile Thr Gln Gly 1160 1155 1165 Thr Pro Ala Leu Pro Gln Thr Gly Ile Pro Thr Glu Ala Leu Val Lys 1170 1175 1180 Gly Ser Ile Ser Arg Met Pro Ile Glu Asp Ser Ser Pro Glu Lys Gly 1190 1195 Arg Glu Glu Ala Ala Ser Lys Gly His Val Ile Tyr Glu Gly Lys Ser 1205 1210 1215 Gly His Ile Leu Ser Tyr Asp Asn Ile Lys Asn Ala Arg Glu Gly Thr 1220 1225 1230 Arg Ser Pro Arg Thr Ala His Glu Ile Ser Leu Lys Arg Ser Tyr Glu 1235 1240 1245 Ser Val Glu Gly Asn Ile Lys Gln Gly Met Ser Met Arg Glu Ser Pro 1260 1255 Val Ser Ala Pro Leu Glu Gly Leu Ile Cys Arg Ala Leu Pro Arg Gly 1270 1275 Ser Pro His Ser Asp Leu Lys Glu Arg Thr Val Leu Ser Gly Ser Ile 1285 1290 Met Gln Gly Thr Pro Arg Ala Thr Thr Glu Ser Phe Glu Asp Gly Leu 1300 1305 1310 Lys Tyr Pro Lys Gln Ile Lys Arg Glu Ser Pro Pro Ile Arg Ala Phe 1315 1320 1325 Glu Gly Ala Ile Thr Lys Gly Lys Pro Tyr Asp Gly Ile Thr Thr Ile 1335 1340 Lys Glu Met Gly Arg Ser Ile His Glu Ile Pro Arg Gln Asp Ile Leu 1350 1355 Thr Gln Glu Ser Arg Lys Thr Pro Glu Val Val Gln Ser Thr Arg Pro 1370 1365 Ile Ile Glu Gly Ser Ile Ser Gln Gly Thr Pro Ile Lys Phe Asp Asn 1380 1385 Asn Ser Gly Gln Ser Ala Ile Lys His Asn Val Lys Ser Leu Ile Thr 1400 1405 Gly Pro Ser Lys Leu Ser Arg Gly Met Pro Pro Leu Glu Ile Val Pro 1415 1420 Glu Asn Ile Lys Val Val Glu Arg Gly Lys Tyr Glu Asp Val Lys Ala 1430 1435 Gly Glu Thr Val Arg Ser Arg His Thr Ser Val Val Ser Ser Gly Pro 1445 1450 Ser Val Leu Arg Ser Thr Leu His Glu Ala Pro Lys Ala Gln Leu Ser 1460 1465 Pro Gly Ile Tyr Asp Asp Thr Ser Ala Arg Arg Thr Pro Val Ser Tyr 1480 1485 Gln Asn Thr Met Ser Arg Gly Ser Pro Met Met Asn Arg Thr Ser Asp 1495 1500 Val Thr Ile Pro Pro Asn Lys Ser Thr Asn His Glu Arg Lys Ser Thr

Leu Thr Pro Thr Gln Arg Glu Ser Ile Pro Ala Lys Ser Pro Val Pro Gly Val Asp Pro Val Val Ser His Ser Pro Phe Asp Pro His His Arg Gly Ser Thr Ala Gly Glu Val Tyr Trp Ser His Leu Pro Thr Gln Leu Asp Pro Ala Met Pro Phe His Arg Ala Leu Asp Pro Ala Ala Ala Tyr Leu Phe Gln Arg Gln Leu Ser Pro Thr Pro Gly Tyr Pro Ser Gln Tyr Gln Leu Tyr Ala Met Glu Asn Thr Arg Gln Thr Ile Leu Asn Asp Tyr Ile Thr Ser Gln Gln Met Gln Val Asn Leu Arg Pro Asp Val Ala 1620 1625 1630 Arg Gly Leu Ser Pro Arg Glu Gln Pro Leu Gly Leu Pro Tyr Pro Ala 1635 1640 Thr Arg Gly Ile Ile Asp Leu Thr Asn Met Pro Pro Thr Ile Leu Val Pro His Pro Gly Gly Thr Ser Thr Pro Pro Met Asp Arg Ile Thr Tyr Ile Pro Gly Thr Gln Ile Thr Phe Pro Pro Arg Pro Tyr Asn Ser Ala Ser Met Ser Pro Gly His Pro Thr His Leu Ala Ala Ala Ser Ala Glu Arg Glu Arg Glu Arg Glu Arg Glu Lys Glu Arg Glu Arg Glu Arg Ile Ala Ala Ala Ser Ser Asp Leu Tyr Leu Arg Pro Gly Ser Glu Gln Pro Gly Arg Pro Gly Ser His Gly Tyr Val Arg Ser Pro Ser Pro Ser Val Arg Thr Gln Glu Thr Met Leu Gln Gln Arg Pro Ser Val Phe Gln Gly Thr Asn Gly Thr Ser Val Ile Thr Pro Leu Asp Pro Thr Ala Gln Leu Arg Ile Met Pro Leu Pro Ala Gly Gly Pro Ser Ile Ser Gln Gly Leu Pro Ala Ser Arg Tyr Asn Thr Ala Ala Asp Ala Leu Ala Ala Leu Val Asp Ala Ala Ala Ser Ala Pro Gln Met Asp Val Ser Lys Thr Lys Glu Ser Lys His Glu Ala Ala Arg Leu Glu Glu Asn Leu Arg Ser Arg Ser Ala Ala Val Ser Glu Gln Gln Gln Leu Glu Gln Lys Thr Leu Glu Val Glu Lys Arg Ser Val Gln Cys Leu Tyr Thr Ser Ser Ala Phe Pro Ser Gly Lys Pro Gln Pro His Ser Ser Val Val Tyr Ser Glu Ala Gly Lys Asp Lys Gly Pro Pro Pro Lys Ser Arg Tyr Glu Glu Glu Leu Arg Thr Arg Gly Lys Thr Thr Ile Thr Ala Ala Asn Phe Ile Asp Val Ile Ile Thr Arg Gln Ile Ala Ser Asp Lys Asp Ala Arg Glu Arg Gly Ser Gln Ser Ser Ser Ser Ser Leu Ser Ser His Arg Tyr Glu Thr Pro Ser Asp Ala Ile Glu Val Ile Ser Pro Ala Ser Ser Pro Ala Pro Pro Gln Glu Lys Leu Gln Thr Tyr Gln Pro Glu Val Val Lys Ala Asn

